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CATCGAAGCGTTATGGCCTGGGAGCCACCGGGGGAGACTCAACATCCAGGCAGGCATTAATGAAAATGACTTTTAC
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CATCTGCATGAGGATGGAGATCTTGGGCTGCCACTGCCGATCCTAATAACTATTATCACCGACGTAATGAGATGA
CCACCACGGATGACCTGGATTTTAAGCACCACAACCTATAAGGAAATGCGCCAGTTGATGAAGGTTGTCAATGAAATG
TGCCCCAATATTACCAGGATTTACAACATTGGCAAAAGCCACCAGGGCCTGAAATTGTATGCGGTAGAGATCTCTGA
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(SEQ ID NO:1)

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ISYASHSTRMLTDARRRVCHTEDFQKEEGTVNGASWHTVAGSLNDFSYLHTNCFELSIYVGC DKYPHESEL
PEEWENNRESLIVFMEQVHRGIKGIVRDLQKGISNAVISVEGVNHDIRTASDGDYWRLLNPG EYVVTAK
AEGFITSTKNCMVGYDMGATRCDFTLTKTNLARIREIMETFGKQPVSLPSRRLKLRGRKRRQRG (SEQ
ID NO:2)

Figure 1

09900518 "070601

underlined = deleted in targeting construct

[] = sequence flanking Neo insert in targeting construct

[AGGCTGTCCCACCCACCATCTGCACCCGCTGCAGCGCCCGCGCCCCCTGTCCC

CGCCGTAGTCGTCATTTGTAGCCCGCCTGCCGCTCCCGGGGACCCGATCCTACCC

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GCCGAGCTCTTCTCGCCTTCAATGCA] TGAAGACCTTAGGGTGGAGGAGCAGGAACAGC

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CCAA [GAGGGAGAAGTTAGTTGCAGAGACGCCTCCACCAG] GTAAAAATAGCAACAGAAA

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CATTACCAAGGAAGGAACCTCTCTTGGCTGAGTGACTGGGTGACATCCTATAAAGTCAT

GGTGAGCAATGACAGCCACACATGGGTTACTGTGAAGAATGGATCTGGCGACATGATATT

TGAAGGAAAACAGTGAGAAGGAGATTCTGTGCTCAATGAGCTGCCAGTCCCCATGGTGGC

CCGCTACATTTCGCATAAACCTCAGTCCTGGTTTGATAACGGGAGCATCTGCATGAGGAT

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CCAGGGCCTGAAATTGTATGCGGTAGAGATCTCTGACCATCCTGGGGAACATGAAGTTGG

TGAGCCCCGAGTTCCACTACATCGCAGGGGGCCACGGCAATGAGGTTCTGGGACGAGA

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CCGCTTGGTGGAGGAGACTCGAATCCACATTCTACCCTCCCTCAATCCTGATGGCTATGA

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CATGGTGC

Figure 2A

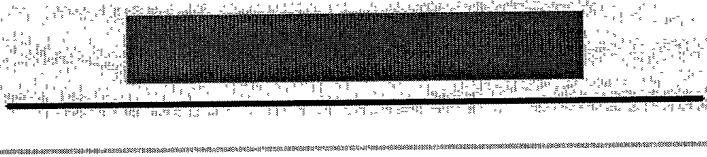
Gene Sequence Structure
*

327 bp

Sequence Deleted

422 bp

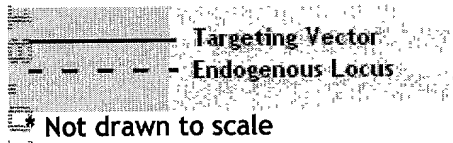
Size of full-length
cDNA: 2490 bp



Targeting Vector* (genomic sequence)

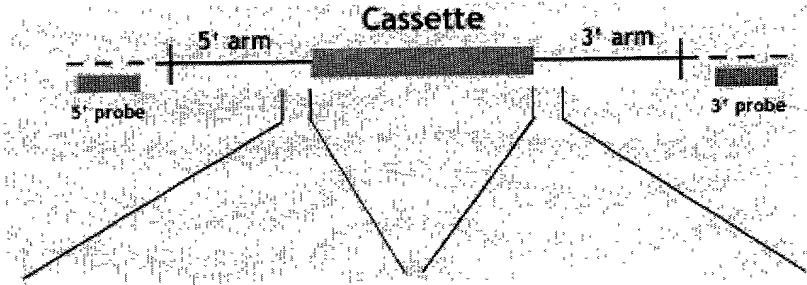
Construct Number: 2825

Arm Length:
5': 4 kb
3': 1.5 kb



LacZ-Neo

Cassette



<p>5' >GGCATGGCCCGTCTG GGGACC GCCTGCCCTGCGCTGGCGCTGGCC CTGGCACTTGTGGCGGTGGCCCTG GCTGGAGTCAGAGCCCAGGGCGCA GCCTTCGAGGAGCCTGACTATTAC AGCCAGGAGCTCTGGCGGCGCGGG CGCTATTATGGGCATCCGGAGCCT GAGCCGGAGCCGGAGCTCTTCTCG CCTTCAATGCA<3' (SEQ ID NO:3)</p>	<p>5' >GAGGGAGAAGTTAGTTGCAGA GACGCCCTCCACCAGGTAAC TTTTG CATCGGGCAGCCCGAGGGGGCGCC AGCGATCGTGGCACTCCAGGGGAC ACCTGGCTTCCAGTATGTTTCTT GAGTGAGCCCAGCCAAAGTCCTGT GGTGCCTGTGTTATTCCCTAGAGA CTACATCTGAGCTAAGTTCAGCTT TCTCTCCCTGC<3' (SEQ ID NO:4)</p>
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Figure 2B

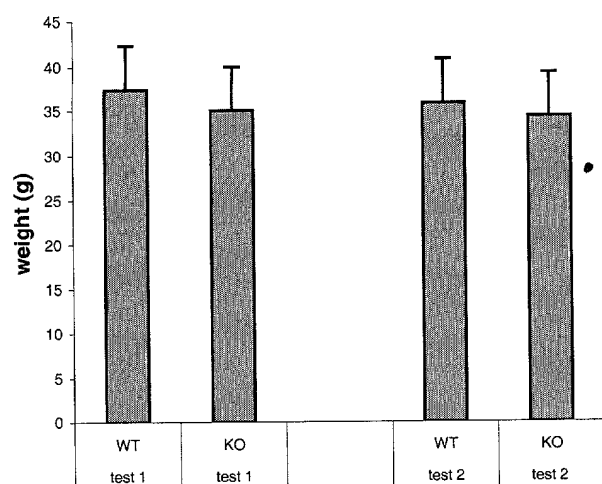


Figure 3

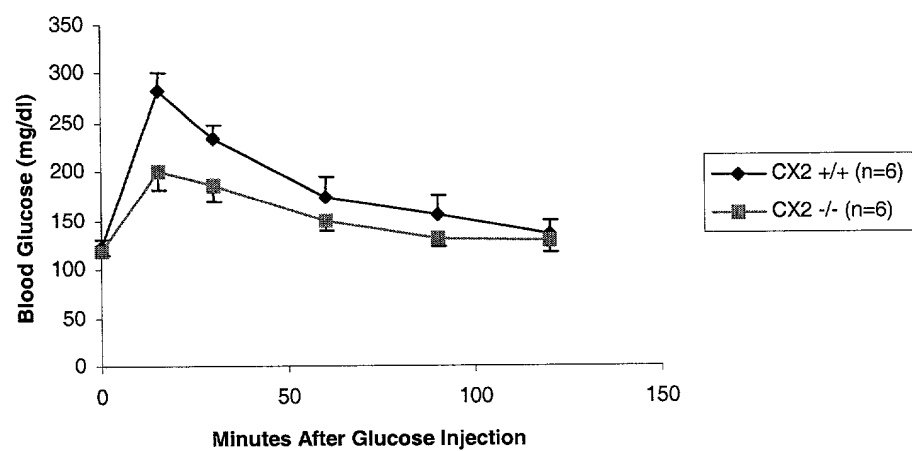


Figure 4